## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 0/5/8, 498Source: PCTDate Processed by STIC: 08/22/2005

## ENTERED



RAW SEQUENCE LISTING

DATE: 08/22/2005

PATENT APPLICATION: US/10/518,498

TIME: 14:11:18

Input Set : A:\PTO.SR.txt

Output Set: N:\CRF4\08222005\J518498.raw

- 3 <110> APPLICANT: november Aktiengesellschaft Gesellschaft for Molekulare Medizin
- 5 <120> TITLE OF INVENTION: Device for detecting an analyte
- 6 <130> FILE REFERENCE: 432739EH
- C--> 8 <140> CURRENT APPLICATION NUMBER: US/10/518,498
- C--> 9 <141> CURRENT FILING DATE: 2004-12-20
  - 11 <160> NUMBER OF SEQ ID NOS: 2
  - 13 <170> SOFTWARE: PatentIn Ver. 2.1
  - 15 <210> SEQ ID NO: 1
  - 16 <211> LENGTH: 20
  - 17 <212> TYPE: DNA
  - 18 <213> ORGANISM: Homo sapiens
  - 20 <220> FEATURE:
- W--> 21 <221> NAME/KEY: modified base
  - 22 <222> LOCATION: (4)
  - 23 <223> OTHER INFORMATION: i
  - 25 <400> SEQUENCE: 1
- W--> 26 cctnccccaa tccctttatt 20
  - 28 <210> SEQ ID NO: 2
  - 29 <211> LENGTH: 20
  - 30 <212> TYPE: DNA
  - 31 <213> ORGANISM: Homo sapiens
  - 33 <400> SEQUENCE: 2
  - 34 aataaaggga ttggggcagg 20



RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/518,498

DATE: 08/22/2005 TIME: 14:11:19

Input Set : A:\PTO.SR.txt

Output Set: N:\CRF4\08222005\J518498.raw

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 4



DATE: 08/22/2005

TIME: 14:11:19



## VERIFICATION SUMMARY

PATENT APPLICATION: US/10/518,498

Input Set : A:\PTO.SR.txt

Output Set: N:\CRF4\08222005\J518498.raw

L:8 M:270 C: Current Application Number differs, Replaced Application Number

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:21 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1

L:26 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0